
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Fri May 25 12:33:28 EDT 2007

Reviewer Comments:

<210> 5

<211> 96

<212> DNA

<213> Artificial sequence

<220>

<223> Fig. 1 A

<400> 5

Invalid <223> Response, if <213> response is Artificial, or unknown please explain the source of genetic material.

Please check for similar errors.

Le A 36 810-Foreign Countries

- 2 -

Please delete this end of file (garble matter in the end of file)

Validated By CRFValidator v 1.0.2

Application No: 10567279 Version No: 1.0

Input Set:

Output Set:

Started: 2007-05-21 11:58:20.621 **Finished:** 2007-05-21 11:58:21.454

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 833 ms

Total Warnings: 10
Total Errors: 6

No. of SeqIDs Defined: 6

Actual SeqID Count: 6

Error code		Error Description
W	213	Artificial or Unknown found in <213> in SEQ ID (1)
W	213	Artificial or Unknown found in <213> in SEQ ID (2)
W	213	Artificial or Unknown found in <213> in SEQ ID (3)
W	213	Artificial or Unknown found in <213> in SEQ ID (4)
W	213	Artificial or Unknown found in <213> in SEQ ID (5)
W	213	Artificial or Unknown found in <213> in SEQ ID (6)
W	112	Upper case found in data; Found at position(104) SEQID(6)
W	112	Upper case found in data; Found at position(106) SEQID(6)
W	112	Upper case found in data; Found at position(113) SEQID(6)
E	342	'n' position not defined found at POS: 120 SEQID(6)
E	342	'n' position not defined found at POS: 120 SEQID(6)
W	112	Upper case found in data; Found at position(120) SEQID(6)
E	342	'n' position not defined found at POS: 124 SEQID(6)
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E	254	The total number of bases conflicts with running total Input: 0, Calculated: 129 SEQID(6)
E	253	The number of bases differs from <211> Input: 104 Calculated:129

SEQUENCE LISTING <110> Bayer AG <120> Screening assay for anti-bacterial compounds <130> LeA36810-EP <140> 10567279 <141> 2007-05-21 <160> 6 <170> PatentIn version 3.1 <210> 1 <211> 26 <212> DNA <213> Artificial sequence <220> <223> Primer T7FF1 <400> 1 gcgcggatcc aaaggaaaat aggagg 26 <210> 2 <211> 30 <212> DNA <213> Artificial sequence <220> <223> Primer T7FF2 <400> 2 30 atcctgaaac tgactgaact aattgagtcg <210> 3 <211> 34 <212> DNA <213> Artificial sequence <220> <223> Primer Xyl1 <400> 3 34 gcgcattaag ctttttctca aggcagtcca attc <210> 4 <211> 96 <212> DNA <213> Artificial sequence <220> <223> Primer Xyl2

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Le A 36 810-Foreign Countries

- 2 -

<210> 5